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	DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI,TDBD; PLUR=YES; OP=OR					
	L30	L29	9			
	DB=PGPB,USPT; PLUR=YES; OP=OR					
	L29	briand-jacques.in.	9			
	L28	L11 and L27	0			
	L27	(enzyme and product) with NMR	118			
	L26	5804390.pn. or 5698401.pn.	2			
	L25	5804390.pn	. 0			
	L24	isotop\$ and 119	1			
	L23	(1H or 3H or 11B or 13C or 15N or 19F or 29S or 31P) and L22	0			
	L22	L21	1			
	L21	chemical adj shift and L19	1			
	L20	chemical ajd shift and L19	898333			
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	L15	WO-9857155-A1.did.	1			
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	L14	L13 not 110	1			
	L13	L11 and L12	5			
	L12	(substrate or ligand) with NMR	1730			
	L11	NMR with (one adj dimension\$ adj spectr\$ or two adj dimension\$ adj spectr\$ or three adj dimension\$ adj spectr\$)	35			
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	L10	L7 and L9	4			
	L9	NMR with (one adj dimension\$ adj spectr\$ or two adj dimension\$ adj spectr\$ or three adj dimension\$ adj spectr\$)	29			
	L8	L5 and L7	238			

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	L7	(substrate or ligand) with NMR	1579	
	L6	12 and L5	355	
	L5	NMR with (one adj dimension\$ or two adj dimension\$ or three adj dimension\$)	2019	
	L4	NMR same (one adj dimension\$ or two adj dimension\$ or three adj dimension\$)	3112	
	L3	11 AND L2	1303	
	L2	(substrate or ligand) same NMR	4300	
	L1	NMR and (one adj dimension\$ or two adj dimension\$ or three adj dimension\$)	12089	

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=> s nmr and (substrate or ligand or enzyme)
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=> s 11 and 12
         49788 L1 AND L2
L3
=> e briand jacques/in
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                 BRIAND J M/IN
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            4 --> BRIAND JACQUES/IN
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            1
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            1
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                 BRIAND L/IN
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E2
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            24 --> BRIAND JACQUES/AU
E3
                   BRIAND JEAN/AU
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                   BRIAND JEAN CLAUDE/AU
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=> s e3
L4
            24 "BRIAND JACQUES"/AU
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FILE 'MEDLINE' ENTERED AT 12:02:06 ON 21 OCT 2004

=> s 14 and 11 3 L4 AND L1 L5

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2 DUP REM L5 (1 DUPLICATE REMOVED)

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Mechanism of Inhibition of Cathepsin K by Potent, Selective ΤI 1,5-Diacylcarbohydrazides: A New Class of Mechanism-Based Inhibitors of Thiol Proteases

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Design of potent and selective human cathepsin K inhibitors that span the TΤ active site.

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ANSWER 1 OF 2 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER:

1999:709526 CAPLUS

DOCUMENT NUMBER:

132:44920

TITLE:

Mechanism of Inhibition of Cathepsin K by Potent, Selective 1,5-Diacylcarbohydrazides: A New Class of

Mechanism-Based Inhibitors of Thiol Proteases

AUTHOR(S):

Bossard, Mary J.; Tomaszek, Thaddeus A.; Levy, Mark

A.; Ijames, Carl F.; Huddleston, Michael J.; Briand, Jacques; Thompson, Scott; Halpert, Stacie; Veber, Daniel F.; Carr, Steven A.; Meek,

Thomas D.; Tew, David G.

CORPORATE SOURCE:

Departments of Molecular Recognition Physical and Structural Chemistry and Medicinal Chemistry,

SmithKline Beecham Pharmaceuticals, King of Prussia,

PA, 19406, USA

SOURCE:

Biochemistry (1999), 38(48), 15893-15902

CODEN: BICHAW; ISSN: 0006-2960

PUBLISHER:

American Chemical Society

DOCUMENT TYPE:

Journal English

LANGUAGE:

The nature of the inhibition of thiol proteases by a new class of AΒ mechanism-based inhibitors, 1,5-diacylcarbohydrazides, is described. These potent, time-dependent, active-site spanning inhibitors include compds. that are selective for cathepsin K, a cysteine protease unique to osteoclasts. The 1,5-diacylcarbohydrazides are slow substrates for members of the papain superfamily with inhibition resulting from slow enzyme decarbamylation. Enzyme-catalyzed hydrolysis of 2,2'-N,N'-bis(benzyloxycarbonyl)-L-leucinylcarbohydrazide is accompanied by formation of a hydrazide-containing product and a carbamyl-enzyme intermediate that is sufficiently stable to be observed by mass spectrometry and NMR. Stopped-flow studies yield a saturation limited value of 43

s-1 for the rate of cathepsin K acylation by 2,2'-N,N'bis(benzyloxycarbonyl)-L-leucinylcarbohydrazide. Inhibition potency varies among proteases tested as reflected by 2-3 orders of magnitude differences in Ki and kobs/I, but all eventually form the same stable covalent intermediate. Reactivation rates are equivalent for all enzymes tested (1 + 10-4 s-1), indicating hydrolysis of a common carbamyl-enzyme form. $\ensuremath{{\bf NMR}}$ spectroscopic studies with cathepsin K and 2,2'-N,N'-bis(benzyloxycarbonyl)-L-leucinylcarbohydrazide provide evidence of inhibitor cleavage to generate a covalent carbamyl-enzyme intermediate rather than a tetrahedral complex. The product Cbz-Leu-hydrazide does not appear enzyme-bound after cleavage in the NMR spectra, suggesting that the stable inhibited form of the enzyme is the thioester complex. 1,5-Diacylcarbohydrazides represent a new class of unreactive cysteine protease inhibitors that share a common mechanism of action across members of the papain superfamily. Both S and S' subsite interactions are exploited in achieving high selectivity and potency.

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DUPLICATE 1

ACCESSION NUMBER:

1998:82491 BIOSIS PREV199800082491

DOCUMENT NUMBER: TITLE:

Design of potent and selective human cathepsin K inhibitors

that span the active site.

AUTHOR(S):

Thompson, Scott K.; Halbert, Stacie M.; Bossard, Mary J.; Tomaszek, Thaddeus A.; Levy, Mark A.; Zhao, Baoguang; Smith, Ward W.; Abdel-Meguid, Sherin S.; Janson, Cheryl A.; D'Alessio, Karla J.; McQueney, Michael S.; Amegadzie, Bernard Y.; Hanning, Charles R.; Desjarlais, Renee L.;

Briand, Jacques; Sarkar, Susanta K.; Huddleston,

Michael J.; Ijames, Carl F.; Carr, Steven A.; Garnes, Keith T.; Shu, Art; Heys, J. Richard; Bradbeer, Jeremy; Zembryki, Denise; Lee-Rykaczewski, Liz; James, Ian E.; Lark, Michael W.; Drake, Fred H.; Gowen, Maxine; Gleason, John G.; Veber,

Daniel F. [Reprint author]

CORPORATE SOURCE:

Dep. Medicinal Chem., SmithKline Beecham Pharm., 709 Swedeland Road, P.O. Box 1539, King Prussia, PA 19406, USA Proceedings of the National Academy of Sciences of the

SOURCE:

Proceedings of the National Academy of Sciences of the United States of America, (Dec. 23, 1997) Vol. 94, No. 26, pp. 14249-14254. print.

CODEN: PNASA6. ISSN: 0027-8424.

DOCUMENT TYPE:

Article English

LANGUAGE: ENTRY DATE:

Entered STN: 24 Feb 1998

Last Updated on STN: 24 Feb 1998

Potent and selective active-site-spanning inhibitors have been designed AΒ for cathepsin K, a cysteine protease unique to osteoclasts. They act by mechanisms that involve tight binding intermediates, potentially on a hydrolytic pathway. X-ray crystallographic, MS, NMR spectroscopic, and kinetic studies of the mechanisms of inhibition indicate that different intermediates or transition states are being represented that are dependent on the conditions of measurement and the specific groups flanking the carbonyl in the inhibitor. The species observed crystallographically are most consistent with tetrahedral intermediates that may be close approximations of those that occur during substrate hydrolysis. Initial kinetic studies suggest the possibility of irreversible and reversible active-site modification. Representative inhibitors have demonstrated antiresorptive activity both in vitro and in vivo and therefore are promising leads for therapeutic agents for the treatment of osteoporosis. Expansion of these inhibitor

concepts can be envisioned for the many other cysteine proteases implicated for therapeutic intervention.

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